

Expression and inheritance of nerve insensitivity resistance in larvae of *Helicoverpa armigera* (Lepidoptera: Noctuidae) from China

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Abstract: Nerve insensitivity resistance to synthetic pyrethroids was detected in a resistant field strain (JSFX-R) of the cotton bollworm, *Helicoverpa armigera* (Hübner), using a neurophysiological assay in which extracellular spontaneous neuronal activity was measured in response to *cis*-cypermethrin. The nerve insensitivity mechanism was selected using a combination of toxicological and neurophysiological methods. The third-instar larvae in selected strains of Family-37 and CTR strain expressed a very high resistance to fenvalerate (RF=2060-fold and 805-fold, respectively) and high cross-resistance to DDT (RF=1927-fold and 2384-fold, respectively) which was not affected by two metabolic synergists, PBO and DMC. The frequency of nerve-insensitive individuals detected in neurophysiological assays (54, 81 and 100% for JSFX-R strain, and the selected strains Family-43 and Family-37, respectively) was not only positively correlated ($R^2=0.968$) with the frequency of non-PBO-synergisable resistant individuals detected in toxicological tests (37.5, 62.5 and 90% for JSFX-R strain, Family-43 and Family-37, respectively), but also positively correlated ($R^2=0.978$) with the frequency of DDT-resistant individuals detected in toxicological tests (40, 67.5 and 93.3% for JSFX-R strain, Family-43 and Family-37, respectively). Analysis of dose-mortality lines to DDT and fenvalerate from F_1 hybrids ($R♀ \times S♂$) indicated that nerve insensitivity resistance to DDT and fenvalerate in the CTR strain was inherited in an incompletely recessive pattern. Degree of dominance (D) was estimated to be $-0.66 (\pm 0.06)$ (DDT) and $-0.26 (\pm 0.04)$ (fenvalerate). The dose-mortality curves to DDT in back-cross progeny were strongly suggested, by chi-square analysis, to be fitted with those expected of a one-gene model. Evidence for the co-existence of nerve insensitivity and oxidative metabolic resistance mechanisms within individual *H. armigera* and the effects of their interaction on the expression of resistance to fenvalerate are discussed.

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1 INTRODUCTION

The cotton bollworm, *Helicoverpa armigera* (Hübner) (previously *Heliothis armigera*), is the main pest of cotton in China and its control has long been a severe problem for Chinese cotton producers. Pyrethroid insecticides such as fenvalerate and deltamethrin were introduced in 1980 and mainly used in cotton for the control of *H. armigera* and the cotton aphid, *Aphis gossypii* (Glov). In 1989 poor control of *H. armigera* by fenvalerate was obtained in the major cotton growing areas of Shandong Province and resistance to fenvalerate was subsequently demonstrated by laboratory bioassays.¹

Resistance to synthetic pyrethroids in *H. armigera* has developed in Asia, the Indian subcontinent, Australia and some African countries,^{2–8} and is threatening the production of cotton and efficacy of pyrethroid insecticides. The mechanisms of resistance

to these compounds have been extensively studied in *H. armigera* and include delayed penetration, enhanced metabolic detoxification and reduced nerve sensitivity.^{9–15} The *kdr*-like nerve insensitivity resistance mechanism has been shown to be an important mechanism in resistance to pyrethroids and DDT in a number of insect pests, on the basis of cross-resistance patterns and an absence of synergism by metabolic synergists.^{16,17} Three possible changes have been thought to account for the reduced sensitivity of nerves to pyrethroids and DDT in *kdr* insects, including structural modifications of the sodium channel leading to reduced receptor binding,^{18–20} alterations in the lipid composition of neuronal membranes,²¹ and a reduction in the density of sodium channels.^{22,23} A number of recent studies have revealed that specific sodium channel mutations may be associated with *kdr*-type nerve insensitivity

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resistance to pyrethroid insecticides.^{24–29} A mutation encoding a leucine to phenylalanine change in transmembrane segment 6 of homology domain II (IIS6) appears to be highly conserved in different insect species^{24–26} although other mutations may also be involved.^{27–29} The existence of this neurophysiologically detectable nerve insensitivity mechanism has been documented in a number of lepidopteran pest species including *Spodoptera littoralis* Boisdu, ³⁰ *Trichoplusia ni* (Hübner),³¹ *Heliothis virescens* F,^{32,33} and *H. armigera* from Thailand,¹⁰ Australia¹¹ and India.¹³ The increased frequency of this mechanism in field populations of the insect has resulted from extensive selection with pyrethroids^{9,11,34} and has led to severe consequences for its control. Early detection is therefore critical to the development of rational strategies for resistance management.

A thorough knowledge of insecticide resistance genetics and the underlying biochemical and physiological mechanisms is useful for devising strategies of insecticide resistance. The detection, monitoring, modelling and risk assessment of resistance can also greatly benefit from a knowledge of its mode of inheritance. The genetic basis of laboratory-selected resistance to pyrethroids has been studied for only a few species of agricultural insect pests, although Georgiou³⁵ reported that resistance to pyrethroids has been documented in at least 48 species of insects. With regard to the mode of inheritance of pyrethroid resistance, no consistent pattern has been observed in the lepidopteran pests which have been reported to date. An incompletely recessive pattern was observed for fenvalerate and permethrin resistance in the diamondback moth, *Plutella xylostella* (L),^{36,37} and for permethrin resistance in the tobacco budworm.³⁸ However, an incompletely dominant pattern was obtained for fenvalerate resistance in the cotton bollworm,³⁹ and for permethrin resistance in the pink bollworm, *Pectinophora gossypiella* (Saunders).⁴⁰ This kind of inconsistency could be due to the different mechanisms in the strain observed or their interactions.

We have previously reported that the presence of the nerve insensitivity mechanism, as indicated by neurophysiological study, is involved in fenvalerate resistance in cotton bollworm from China.⁴¹ Herein we report the expression of the nerve insensitivity resistance to DDT and fenvalerate in selected strains of *H. armigera* and the inheritance pattern of this mechanism in a selected homozygous (CTR) strain.

2 MATERIALS AND METHODS

2.1 Insects

All the studies were performed using third-instar larvae of various strains of *Helicoverpa armigera*. A laboratory strain, Reading, was used as a susceptible reference strain and had been kept in culture for over 14 years without selection. The JSFX-R strain originated from samples of at least 1000 eggs and

neonate larvae of a pyrethroid-resistant population of *H. armigera*, collected randomly from cotton plants in the Fengxie county area of Jiangsu province in China in June 1994 and June–July 1995. From this strain, further resistant strains were obtained by selective breeding as described below.

2.2 Insecticides

Technical *cis*-cypermethrin (98.4%) and fenvalerate (97.5%) were supplied by Zeneca Agrochemicals, Jealott's Hill, UK and Shell Research Limited, Sittingbourne, UK, respectively. Piperonyl butoxide (2-(2-butoxyethoxy)ethyl 6-propylpiperonyl ether; PBO) was supplied as technical material (98%) by Endura, Italy. 1,1-bis(4-Chlorophenyl)-2,2,2-trichloroethane (DDT) and 1,1-bis(4-chlorophenyl)-ethanol (DMC) were supplied by Aldrich Chemical Company Ltd, Gillingham, UK, as technical material (98% and 97% respectively).

2.3 Neurophysiological assay for nerve insensitivity

Extracellular spontaneous neuronal activity was recorded in the exposed peripheral nerves of partly dissected third-instar larvae bathed in saline containing *cis*-cypermethrin at 25(±2)°C using a cumulative dose response assay described in detail previously by McCaffery *et al.*^{33,41} EC₅₀ values were estimated using a probit analysis. Highly resistant insects not responding at 100 nM *cis*-cypermethrin could not be included in this estimation. In Family-37 EC₅₀ was estimated as >100 nM because all tested insects responded at or above the highest concentration of 100 nM *cis*-cypermethrin.

2.4 Insecticide and synergist bioassays

The toxicity of fenvalerate and DDT to the third-instar larvae with a body weight of 13–20 mg in all the strains of *H. armigera* was determined in full dose mortality topical bioassays as described previously.^{4,42} Insecticide was dissolved in analytical grade acetone and appropriate serial dilution were prepared. A fixed volume of insecticide solution (0.2 µl) was topically applied on the thoracic dorsum of each third-instar larva using a Hamilton syringe dispenser. In the case of very high concentrations of DDT, where crystallisation could occur, the solution was diluted and a 1-µl drop of insecticide solution was applied on each larva. Newly moulted larvae or those about to moult were excluded from all tests. Three or four replicates of 10 larvae were treated for the control and each concentration and at least four or five concentrations in the appropriate range were used for each test. All treated insects were kept separately in 30-ml sterile pots with a small block of artificial diet, under the same conditions as the stock insect cultures. Mortality was assessed 72 h after treatment. Larvae were classified as dead if they were unable to change positions within 15 s after being prodded with a blunt seeker. Synergist bioassays were performed by pre-dosing test insects

with 20 µg synergist (PBO or DMC) in 0.2 µl of acetone on the dorsal portion of the abdomen 30 min prior to the application of a range of insecticide doses. Control insects were treated with acetone alone or treated with synergist alone. No mortality was found in control insects treated with acetone or with synergist alone. Toxicity was evaluated using SAS software⁴³ and Micro-Probit 3.0

2.5 Selection of nerve insensitivity resistance mechanism

In order to form a homozygous strain expressing the nerve insensitivity resistance mechanism, the JSFX-R strain was selected and purified using a combination of single pair mating, toxicological and neurophysiological methods. Insects of the JSFX-R strain from F₂ to F₄ were selected in the third stadium using a diagnostic dose of fenvalerate of 0.04 µg per larva. The offspring of single pair matings at F₅ were screened with a diagnostic dose of fenvalerate following pre-treatment (30 min) with 20 µg PBO. Families which expressed lower mortality to fenvalerate alone, together with lower synergism by PBO, were kept for further selection. Resistant Family-43 at F₁₀ was chosen as the line for further selection based on a high resistance to fenvalerate and a low PBO synergistic ratio. Family-37 at F₁₄, which showed high resistance to fenvalerate and which was homozygous for nerve insensitivity, as indicated by the neurophysiological assay, was kept as the nerve-insensitive strain. Between the generations where single pair matings were set up, all third-instar larvae were selected with a diagnostic dose of fenvalerate. Finally, the CTR strain was derived from the offspring of Family-37, which was selected with fenvalerate.

2.6 Inheritance study

The Reading strain was used as the susceptible parent and the CTR strain was used as the resistant parent. Virgin females and males were obtained by determining the sex of pupae. Female and male pupae were held in separate dishes to await adult emergence. Reciprocal F₁ crosses (S♂ × R♀ and S♀ × R♂) were made by mass matings between Reading strain (S) and CTR strain (R). A portion of the F₁ progeny was used in the assessment of dose–mortality response. Others were reared to maturity for F₂ crosses which include F₁♂ × F₁♀ and F₁♂ × R♀. The dose–mortality response in all crosses was determined using the topical bioassay methods described above.

2.7 Data analyses

The dose–mortality relationship obtained from F₁ cross and from parents was used to determine the degree of dominance (*D* value) of fenvalerate and DDT resistance using the formula

$$D = \frac{2X_3 - X_2 - X_1}{X_2 - X_1}$$

where X_1 , X_2 and X_3 were the log₁₀(LD₅₀) of susceptible Reading strain, resistant CTR strain and heterozygous F₁ hybrid.⁴⁴ The standard error of the dominance value (*D*) was calculated by taking the square root of variance of *D* value⁴⁵

$$\text{var}(D) = \frac{4}{(X_2 - X_1)^2}$$

$$\cdot \left[\text{var}(X_3) + \frac{(X_3 - X_1)^2}{(X_2 - X_1)^2} \cdot \text{var}(X_2) + \frac{(X_3 - X_2)^2}{(X_2 - X_1)^2} \cdot \text{var}(X_1) \right]$$

The standard error (SE) of a *D* value is useful in determining whether the value of *D* is significantly different from ±1 or 0. Statistical analysis of back-cross data was carried out with a chi-square test for the goodness-fit between the observed response and expected response at each dose in a series, as described by Preisler *et al.*⁴⁵

$$\chi^2 = \frac{(r_i - n_i p_i)^2}{n_i p_i (1 - p_i)}$$

where n_i = total number of insects from the back-cross generation treated with the *i*th dose. r_i = observed number of dead at *i*th dose and p_i = estimated mortality probability under the hypothesised genetic model at *i*th dose. Expected mortalities were estimated assuming that resistance to DDT or fenvalerate is determined by a single major gene with a Mendelian mode of inheritance.⁴⁶ Therefore expected mortality at each dose for the back-cross of F₁♂ × R♀ and inter-cross of F₁♂ × F₁♀ was calculated respectively as:

$$X_i = 0.5 W_{F1} + 0.5 W_{RR}$$

$$X_i = 0.25 W_{SS} + 0.5 W_{F1} + 0.25 W_{RR}$$

where W_{F1} is the observed mortality at *i*th dose of the F₁ hybrid (S♂ × R♀), W_{RR} is the observed mortality at *i*th dose of resistant homozygous parent CTR strain, and W_{SS} is observed mortality at *i*th dose of susceptible homozygous parent Reading strain. The genetic hypothesis was then tested by comparing the test static χ_i^2 (for each $i = 1 \dots N$) with values from χ^2 table with 1 df, or by comparing the sum $\chi^2 = \sum_i \chi_i^2$ with values from χ^2 table with *N* df.

3 RESULTS

3.1 Resistance to fenvalerate and DDT and synergism

3.1.1 Resistance to fenvalerate

A full dose–mortality assay with fenvalerate was conducted using the Reading susceptible strain, the JSFX-R field strain, and the selected CTR strain. A high level of resistance to fenvalerate (RF = 200-fold at the LD₅₀ value) was observed in the JSFX-R strain relative to the Reading strain. After selection, the resistance to fenvalerate increased to 2060-fold in Family-37 and 805-fold in the CTR strain (Table 1). Pre-treatment of the Reading susceptible strain with

Table 1. Toxicity of fenvalerate to third-instar larvae of the Reading susceptible and various resistant strains of *Helicoverpa armigera* and synergism with piperonyl butoxide (PBO)

Strain	Treatment	LD ₅₀ (µg per larva) (95% fiducial limits)	RF ^a	SR ^b	Slope (± SE)
Reading	Fenvalerate	0.003 (0.0027~0.0033)	1.0		2.9 (±0.2)
Reading	Fenvalerate + PBO	0.008 (0.0063~0.0095)	2.7	0.4	2.4 (±0.3)
JSFX-R	Fenvalerate	0.60 (0.40~1.09)	200		1.6 (±0.3)
JSFX-R	Fenvalerate + PBO	0.018 (0.013~0.027)	6.0	33.5	1.4 (±0.2)
Family-37	Fenvalerate	6.18 (4.17~8.91)	2060		1.1 (±0.1)
Family-37	Fenvalerate + PBO	0.14 (0.11~0.18)	46.7	44.1	1.8 (±0.2)
CTR	Fenvalerate	1.69 (1.25~2.53)	805		1.3 (±0.1)

^a Resistance factor (RF) expressed as LD₅₀ of resistant strain/LD₅₀ of Reading strain.^b Synergistic ratio (SR) expressed as LD₅₀ without PBO/LD₅₀ with PBO.**Table 2.** Toxicity of DDT to third-instar larvae of susceptible Reading strain and resistant Family-37 of *Helicoverpa armigera* and synergism with PBO and DMC

Treatment	Reading strain			Family-37			
	LD ₅₀ (µg per larva)	Slope (± SE)	SR ^b	LD ₅₀ (µg per larva)	Slope (± SE)	SR ^b	RF ^a
DDT	0.26	2.16 (±0.41)		501	0.93 (±0.18)		1927
DDT + PBO	0.43	2.94 (±0.47)	0.60	558	0.92 (±0.20)	0.90	1298
DDT + DMC	0.39	2.04 (±0.40)	0.67	548	0.72 (±0.19)	0.91	1405

^a Resistance factor (RF) expressed as LD₅₀ of resistant Family-37/LD₅₀ of Reading strain.^b Synergistic ratio (SR) expressed as LD₅₀ without synergist/LD₅₀ with synergist.**Table 3.** The nerve insensitivity level in different strains, and their toxicological response to fenvalerate and DDT

Strain	Neurophysiology		Toxicology		
	Nerve insensitivity (%) (n)	EC ₅₀ (nM) (95% fiducial limits)	Resistance to fenvalerate (%)	Resistance to fenvalerate/PBO (%)	Resistance to DDT (%)
Reading	0 (44)	1.2 (0.7~1.8)	0	0	0
JSFX-R	54 (26)	12.7 (7.9~20.6)	97.5	37.5	40.0
Family-43	81 (26)	39.9 (26.7~59.8)	73.3	62.5	67.5
Family-37	100 (49)	>100	100	90.0	93.3

Nerve-insensitive individuals were those responding to *cis*-cypermethrin at or above a concentration of 100 nM in the neurophysiological assay. Reading strain individuals all responded at or below 50 nM *cis*-cypermethrin. The values in toxicological tests represent the percentage of resistant individuals surviving the discriminating dose of fenvalerate (0.04 µg per larva), fenvalerate/PBO (0.04/20 µg) and DDT (8 µg per larva).

PBO resulted in no synergism of fenvalerate, but pre-treatment of the JSFX-R strain and Family-37 gave 33.5-fold and 44.1-fold synergism to fenvalerate, respectively (Table 1). The non-PBO-synergisable resistance increased from 6.0-fold in the JSFX-R strain to 46.7-fold in Family-37. This suggested that the nerve insensitivity resistance mechanism was preferentially selected in this process.

3.1.2 Cross-resistance to DDT

A very strong cross-resistance to DDT was observed in larvae from the fenvalerate-resistant Family-37 and CTR strain. The resistance to DDT was over 1927-fold and 2384-fold (at the LD₅₀ value) relative to the Reading susceptible strain (see Tables 2 and 4). This resistance was not suppressed by pre-treatment of larvae with DMC (a DDT dehydrochlorinase inhibitor) or PBO (a monooxygenase inhibitor) (Table 2),

providing further indirect evidence for the presence of nerve insensitivity resistance in these strains.

3.1.3 Correlation between neurophysiological and toxicological responses

The results shown in Table 3 and Fig 1a illustrate that there was a significant positive correlation ($R^2 = 0.968$) between the frequency of nerve-insensitive individuals, detected in neurophysiological assays, and the frequency of non-PBO-synergisable individuals detected in toxicological tests. In the JSFX-R strain the non-PBO-synergisable individuals comprised 37.5% of the population, while in Family-43 and Family-37 the non-PBO-synergisable individuals comprised 67.5 and 90% of the population, respectively (Table 3). The presence of the nerve insensitivity resistance mechanism was again demonstrated by the positive correlation ($R^2 = 0.978$) between the frequency of DDT-resistant individuals

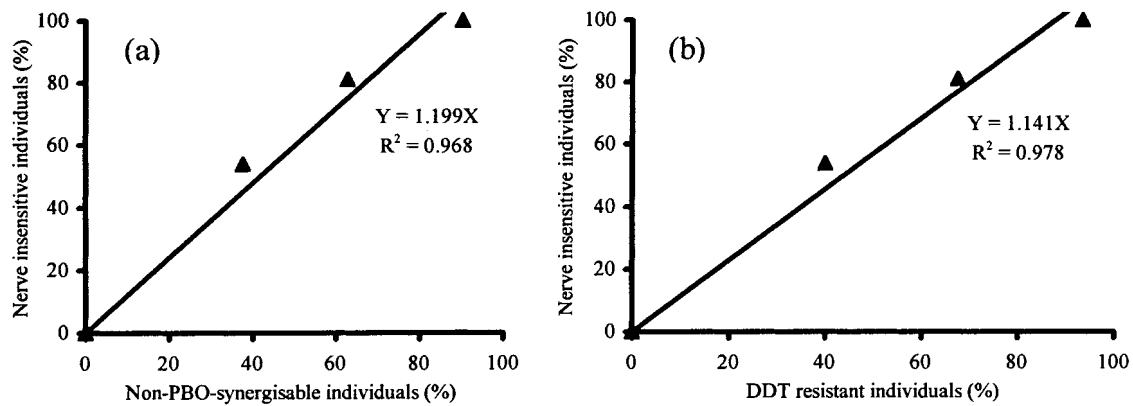


Figure 1. Correlation between (a) non-PBO-synergisable individuals detected in toxicological tests and nerve-insensitive individuals detected in neurophysiological assays and (b) DDT-resistant individuals detected in toxicological tests and nerve-insensitive individuals detected in neurophysiological assays. Significance level $P < 0.05$.

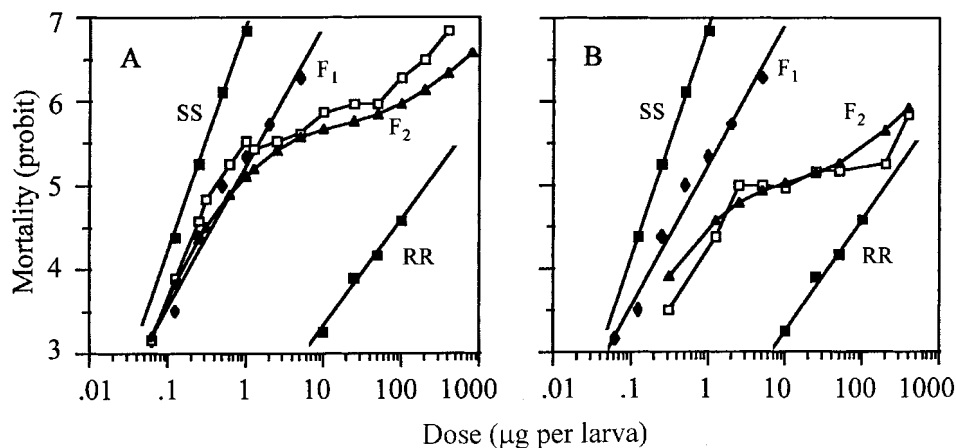


Figure 2. Dose-mortality responses for genetic crosses of nerve insensitivity resistant CTR strain of *Helicoverpa armigera* when tested with DDT. The solid line (\blacktriangle) shows the expected response for (A) a 1:2:1 segregation in the inter-cross of $F_1\delta \times F_1\varphi$ and for (B) a 1:1 segregation in the backcross of $F_1\delta \times R\varphi$. The expected mortalities at each dose are estimates based on simple Mendelian inheritance of one-locus model (Georghiou, Reference 46).

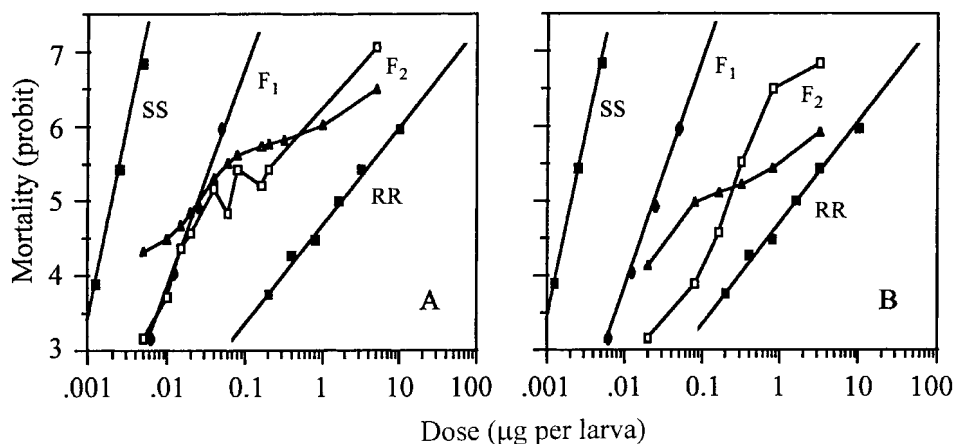


Figure 3. Dose-mortality responses for genetic crosses of nerve insensitivity resistant CTR strain of *Helicoverpa armigera* when tested with fenvalerate. The solid line (\blacktriangle) shows the expected response for (A) a 1:2:1 segregation in the inter-cross of $F_1\delta \times F_1\varphi$ and for (B) a 1:1 segregation in the backcross of $F_1\delta \times R\varphi$. The expected mortalities at each dose are estimates based on simple Mendelian inheritance of one-locus model (Georghiou, Reference 46).

shown by the percentage of insects surviving the twice discriminating dose of DDT ($8\mu\text{g}$ per larva) in toxicological tests and the frequency of nerve-insensitive individuals in neurophysiological assays (Fig 1b).

In the JSFX-R strain the DDT-resistant individuals comprised 40% of the population, and this value increased to 67.5% in Family-43 and 93.3% in Family-37 (Table 3).

Table 4. Responses to fenvalerate and DDT of the third-instar larvae of *Helicoverpa armigera* from the homozygous susceptible Reading strain (SS) and resistant CTR strain (RR), and the F₁ (SS♂ × RR♀)

Strain or cross	n	LD ₅₀ (μg per larva) (95% fiducial limits)	RF ^a	Slope (±SE)	D (±SE) ^b
<i>Fenvalerate</i>					
Reading (SS)	120	0.0021 (0.0018~0.0024)		4.96 (±0.87)	
CTR (RR)	460	1.69 (1.25~2.53)	805	1.34 (±0.13)	
F ₁ (S♂ × R♀)	120	0.067 (0.050~0.094)	31.9	3.14 (±0.51)	−0.26 (±0.04)
<i>DDT</i>					
Reading (SS)	120	0.21 (0.15~0.26)		2.80 (±0.50)	
CTR (RR)	200	501 (253~1912)	2384	0.93 (±0.18)	
F ₁ (S♂ × R♀)	210	0.67 (0.51~0.91)	3.2	1.72 (±0.20)	−0.66 (±0.06)

^a Resistance factor (RF) was determined by dividing the LD₅₀ of resistant strain by the LD₅₀ of susceptible Reading strain.

^b The value of dominance D = −1, completely recessive; D = 0, intermediate or co-dominant; D = +1, completely dominant.

3.2 Inheritance

3.2.1 F₁ dose-response

Responses to DDT and fenvalerate of the susceptible Reading strain, the resistant CTR strain and their crosses are given in Table 4 and Figs 2 and 3. Resistances to DDT and fenvalerate were 2384-fold and 805-fold, respectively, in the CTR strain and 3.2-fold and 11.9-fold, respectively, in the progenies of cross (S♂ × R♀). Although the maternal influence on the resistance cannot be evaluated, the degree of dominance and its standard error were −0.66 (±0.06) for DDT resistance and −0.26 (±0.04) for fenvalerate resistance. These results clearly indicated that resistance to DDT and to fenvalerate in the CTR strain was inherited in an incompletely recessive manner.

3.2.2 F₂ dose-response

The dose-mortality curves to DDT displayed an obvious plateau at 50% mortality in the offspring of F₁ back-cross to resistant parent CTR strain, and a plateau at 75% mortality in the progenies of F₁ inter-cross (Fig 2). Chi-square analysis indicated that the observed dose-mortality curves to DDT in F₂ of F₁♂ × R♀ ($\chi^2_{(0.05)[14]} = 13.08$) and F₁♂ × F₁♀ ($\chi^2_{(0.05)[9]} = 7.94$) were respectively fitted to the expected curves calculated from a monogenic model (Fig 2). Together with the lack of synergism by PBO and DMC in toxicological tests and the observed neurophysiological response, these results showed clearly that resistance to DDT in CTR strain is entirely due to the presence of a mechanism of nerve insensitivity. In contrast, the dose-mortality curves to fenvalerate, in F₂ progenies (Fig 3) were different from the curves expected of a one-gene model. It therefore appears that more than one mechanism was involved in the resistance to fenvalerate, as illustrated by the expression of PBO synergism to fenvalerate in the strain (Table 1).

4 DISCUSSION

The results presented here strongly suggest that a nerve insensitivity resistance mechanism plays an

important role in the resistance to fenvalerate and DDT in the JSFX-R strain of the cotton bollworm collected from China and in the strains derived from the JSFX-R strain. Individuals expressing this mechanism were selected in these studies and, as a result, a strain which was homozygous for the mechanism was obtained. Previous neurophysiological work has concluded that nerve insensitivity is an important mechanism of resistance to pyrethroids in Thai and Indian strains of *H. armigera*.^{10,13} In a study of nerve insensitivity in *H. armigera* in Australia, the time to burst discharge to a high dose of fenvalerate (10^{−5} M) was significantly different between resistant and susceptible insects and could be used to classify the resistant and susceptible insects.¹¹ Together these studies have demonstrated the widespread nature of a nerve insensitivity resistance mechanism which confers resistance to pyrethroid insecticides in *H. armigera*.⁴¹

The expression of this nerve insensitivity resistance mechanism usually confers cross-resistance to DDT and pyrethroids, despite their different chemical structures,⁴⁷ because DDT and pyrethroids are known to exert their toxic effects by binding to and modifying the normal function of voltage-sensitive sodium channels in a very similar way.^{48,49} The appearance of cross-resistance to DDT in a strain resistant to pyrethroids has, therefore, very often been taken as an indication that the nerve insensitivity mechanism is involved in the resistance. Our results (Table 2) indicated that in Family-37 and the CTR strain, which was shown to be homozygous for nerve insensitivity, there is a high level of cross-resistance to DDT without any synergistic action by PBO and DMC. The phenomenon of cross-resistance between pyrethroids and DDT has been well documented in other lepidopteran species such as *Heliothis virescens*,^{50,51} *Phyllonorycter blancardella* F,⁵² *Plutella xylostella*,⁵³ and has been generally thought to be due to the existence of a nerve insensitivity resistance mechanism.^{17,54} The positive correlation ($R^2 = 0.978$) between the percentages of DDT-resistant individuals and nerve-insensitive individuals (Fig 1b) provides

more reliable evidence in support of the presence of cross-resistance caused by the nerve insensitivity mechanism.

Further evidence for the involvement of the nerve insensitivity mechanism is the absence of synergism of pyrethroids or DDT by metabolic synergists such as PBO, DEF (*S,S,S*-tributyl phosphorotrithioate), and DMC.⁵⁵ Liu *et al*⁵³ found resistance to DDT and pyrethroids was unaffected by metabolic synergists in the diamondback moth, *P xylostella*, suggesting that a *kdr* factor could be a resistance mechanism. In studies on mechanisms of resistance to pyrethroids in *H virescens*, non-PBO-synergisable resistance was thought to be due to the nerve insensitivity mechanism.⁵⁶ A significant correlation was found between the numbers of non-PBO-synergisable resistant individuals and the numbers of nerve-insensitive individuals in *H virescens*.³³ The results presented here also indicate a positive correlation between the frequency of nerve-insensitive individuals and non-PBO-synergisable individuals in the fenvalerate-resistant JSFX-R strain and selected strains of *H armigera* ($R^2 = 0.968$). The frequency of non-PBO-synergisable individuals or the frequency of DDT-resistant individuals detected in a resistant field population of *H armigera* could therefore indicate the importance of nerve insensitivity resistance mechanism in a population.

Although Family-37 was confirmed by neurophysiological experiments to be a strain homozygous for nerve insensitivity, the resistance to fenvalerate could still be suppressed to a moderate degree by PBO (Table 1). The contribution of each resistance mechanism to the overall fenvalerate resistance (2060-fold) was 44.1-fold (48.6%) for the non-PBO-synergisable factor, presumed to be largely the nerve insensitivity resistance mechanism, and 46.7-fold (51.4%) for the PBO-synergisable resistance mechanism, if interaction of the two resistance factors in combination was multiplicative.⁵⁷ These results imply that the oxidative metabolic resistance mechanism can coexist with the nerve insensitivity resistance mechanism in individual larvae, and could explain the interaction of these two resistance mechanisms on the expression of resistance to pyrethroids. In a field population of *H armigera* in Australia, pyrethroid selection pressure in the field was found clearly to select for the oxidative metabolic resistance mechanism.⁷ Similarly, in this laboratory, mass selection of third-instar larvae of the JSFX-R strain with fenvalerate by topical application led to a dramatic increase in the level of fenvalerate resistance which could be suppressed by PBO (data not shown). This is likely to be due to the differential genetic dominance of each resistance mechanism. The oxidative metabolic resistance mechanism in Australian *H armigera* has been shown to be semi-dominant,³⁹ whilst the nerve insensitivity resistance mechanism has been considered to be recessive in other insects.^{54,58} The obviously different fitness costs associated with these two resistance mechanisms may have also contributed to the different selection.⁵⁹ In field

populations it is possible that the oxidative metabolic resistance mechanism may protect the nerve insensitivity resistance mechanism, when these two mechanisms coexist in the same individual.

The genetic studies showed that resistance to DDT and fenvalerate in the CTR strain was inherited in an incompletely recessive manner and demonstrated a general characteristic of the nerve insensitivity mechanism.^{54,58} The dose-mortality response to DDT in the back-crosses (Fig 2), relative to the expected dose-response based on a one-gene model, strongly suggested that nerve insensitivity was the major mechanism conferring resistance to DDT in the CTR strain, whilst the dose-mortality response to fenvalerate in the back-crosses (Fig 3) indicated more than one mechanism contributed to resistance to fenvalerate in the CTR strain. Interestingly, the differential of dose-mortality response to DDT and to fenvalerate in the F_2 progeny suggested an alternative way to study the genetics of this specific resistance mechanism in a strain, although there might be more than one mechanism included. Because DDT resistance in the CTR strain was not affected by metabolic synergists such as PBO and DMC (Table 2), the dose-mortality response to DDT could be used as the phenotypic expression of nerve insensitivity in a study of the genetics of this mechanism. Resistance to fenvalerate in the CTR strain involved the co-existence of nerve insensitivity and oxidative metabolism. To test for linkage and the possible interaction of the nerve insensitivity gene with oxidative metabolism, the two factors must first be isolated in isogenic strains. This is presently being undertaken in this laboratory using a series of crosses and back-crosses, together with single-pair mating.

The main aim of insecticide resistance management (IRM) is to conserve or recover susceptibility to an insecticide in a pest population using modified insecticide use patterns which reduce selection.⁶⁰ The results of this study clearly showed that the substantial resistance to fenvalerate in the field resistant strain (JSFX) of *H armigera* from China could be eliminated by low doses of fenvalerate following PBO pre-treatment. This implies that metabolic detoxification, probably via mono-oxygenases, was the major mechanism of resistance in this population. In the field, therefore, a mixture of pyrethroid plus PBO could be used as a strategy for resistance management and to retain susceptibility to the pyrethroids. Although selection with a mixture of a pyrethroid and PBO could lead to the development of a nerve insensitive resistance mechanism, this should be more easily managed due to its genetically recessive nature and associated higher fitness costs. Nevertheless, care would need to be taken to avoid the emergence of a *super-kdr* type of nerve insensitivity resistance mechanism with metabolic resistance mechanisms which would lead to the development of a highly resistant population.^{61,62}

The identification of nerve insensitive and metabolic

resistance to pyrethroids in Chinese strains of *H. armigera* leads to more fundamental considerations of the design of management strategy to control resistance. Resistance to endosulfan, OPs and carbamates arises through mechanisms usually entirely distinct from these identified here in resistance to pyrethroids. A management strategy which employs rotation with endosulfan, OPs and certain carbamates could thus prove useful in reducing the frequency of pyrethroid resistance genes in field populations. Likewise, the use of unsprayed refugia might also prove of enormous benefit to the survival of susceptible alleles and we believe that consideration of these approaches is now appropriate.

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